

XX CC The invention relates to a novel human G-coupled receptor (I). (I) and
CC its corresponding polynucleotides are useful for diagnosing, treating or
CC preventing cell proliferative diseases (e.g. lymphoma, leukaemia, breast
CC cancer or cirrhosis), neurological disorders (e.g. stroke, Alzheimer's
CC disease, multiple sclerosis or mental retardation), cardiovascular
CC diseases (e.g. atherosclerosis, angina pectoris or congestive heart
CC failure), gastrointestinal disorders (e.g. dysphagia, indigestion or
CC gastritis), autoimmune/inflammatory disorders (e.g. AIDS, Crohn's disease
CC or systemic lupus erythematosus) or metabolic disorders (e.g. diabetes or
CC obesity), or viral infections (e.g. infection by herpesvirus or
CC parvovirus). AAU80493-AAU80515 represent novel human G-coupled receptor
CC amino acid sequences of the invention
XX XX
XX Sequence 321 AA;
Query Match 96.0%; Score 1649; DB 5; Length 321;
Best Local Similarity 98.8%; Pred. No. 2.8e-173;
Matches 317; Conservative 1; Mismatches 3; Indels 0; Gaps 0;
QY 9 MESPHTDVPSPVFLGIPGLEQFHLWLSLPVCGLTATVGNITILVVATEPVLHKP 68
Db 1 MESPHTDVPSPVFLGIPGLEQFHLWLSLPVCGLTATVGNITILVVATEPVLHKP 60
QY 69 VYFLCMLSTIDLAASVSTVPKLLAIFWCGAGHISASACLAHMFPIHAFCMESTVLLAM 128
Db 61 VYFLCMLSTIDLAASVSTVPKLLAIFWCGAGHISASACLAHMFPIHAFCMESTVLLAM 120
QY 129 AFDYVAICHPLRYATILDTTIIAHIGVAAVVRGSLMLPCPFFIGRLNFCQSHVILTY 188
Db 121 AFDYVAICHPLRYATILDTTIIAHIGVAAVVRGSLMLPCPFFIGRLNFCQSHVILTY 180
QY 189 CEHMAVVKLAGDTRPNRVYGLTAALVIGVDLFCIGLSYALIAQAVRLSSHEARSKAL 248
Db 181 CEHMAVVKLAGDTRPNRVYGLTAALVIGVDLFCIGLSYALIAQAVRLSSHEARSKAL 240
QY 249 GTCGSHVCVILISYTPALFSPFTHRGHHVPHIHLANVYLLPPALNPVYGVKTKQ 308
Db 241 GTCGSHVCVILISYTPALFSPFTHRGHHVPHIHLANVYLLPPALNPVYGVKTKQ 300
QY 309 IRKRVVRVFGSGQGMGIKASE 329
Db 301 IRKRVVRVFGSGQGMGIKASE 321
RESULT 10
AAU95725
ID AAU95725 standard; protein; 321 AA.
XX AC AAU95725;
XX AC AAU95725;
DT 02-JUL-2002 (first entry)
XX DE Human olfactory and pheromone G protein-coupled receptor #212.
XX KW Human; olfactory and pheromone G protein coupled; receptor; GPCR;
KW tranquilizer; antidepressant; neuroleptic; endocrine; anabolic;
KW anorectic; taste; fragrance; food additive; cosmetic; cell migration;
KW sterility; psychotic disorder; neurological disorder; anxiety;
KW schizophrenia; manic depression; depression; axonal growth;
KW menstrual cycle; appetite sexual motivation; sexual attraction;
KW aggression.
XX OS Homo sapiens.
XX FN WO200224726-A2.
XX FN 28-MAR-2002.
XX PD 21-SEP-2001; 2001WO-BE000162.
XX PF 22-SEP-2000; 2000EP-00870211.
XX PR 22-SEP-2000; 2000EP-00870211.
XX XX

(CHEM-) CHEMCOM SA.
XX Veithen A;
XX WPI; 2002-330013/36.
DR N-PSDB; ABK68612.
XX Novel pheromone G-protein coupled receptor and receptor-derived agonists,
PT antagonists or inhibitors useful in food or cosmetic products or in the
PT treatment or prevention of neurological disorders such as anxiety and
XX schizophrenia.
PS Disclosure; Page 647-648; 833pp; English.
XX The invention relates to olfactory and Pheromone G-protein coupled
CC receptor (GPCR) or a protein 95% identical to the GPCR, a specific active
CC portion and its encoding polynucleotide. Also included are an agonist,
CC antagonist or inhibitor of the GPCR or the polynucleotide, a vector,
CC comprising the polynucleotide, a cell transformed by the vector, a non-
CC human mammal comprising a partial or total deletion of the polynucleotide
CC encoding the receptor and screening (detection and possibly, recovering)
CC of compounds which are known or not known to be agonist, antagonists or
CC inhibitors of natural compounds to the GPCR. The receptor-derived
CC agonists, antagonists, inhibitors or compounds are used as an
CC improvement, elimination or substitution of an existing taste and/or a
CC fragrance of (or in) the food and/or cosmetic products. They can also be
CC used in the preparation of medicament in the treatment and/or prevention
CC of a mammalian disorder, such as cell migration, sterility, psychotic and
CC neurological disorders, including anxiety, schizophrenia, manic
CC depression, depression, for promoting axonal growth, nerve cell
CC connection and nerve regeneration for modulating male and female
CC endocrine functions, hormone production and the menstrual cycle, for the
CC prevention or the treatment by stimulation of several mammalian
CC behaviours, such as stimulation or suppression of appetite, sexual
CC motivation, sexual attraction, aggression and for promoting or
CC suppressing chemical communication between organisms. The present
CC sequence is a human olfactory and pheromone GPCR protein sequence
XX Sequence 321 AA;
Query Match 96.0%; Score 1649; DB 5; Length 321;
Best Local Similarity 98.8%; Pred. No. 2.8e-173;
Matches 317; Conservative 1; Mismatches 3; Indels 0; Gaps 0;
QY 9 MESPHTDVPSPVFLGIPGLEQFHLWLSLPVCGLTATVGNITILVVATEPVLHKP 68
Db 1 MESPHTDVPSPVFLGIPGLEQFHLWLSLPVCGLTATVGNITILVVATEPVLHKP 60
QY 69 VYFLCMLSTIDLAASVSTVPKLLAIFWCGAGHISASACLAHMFPIHAFCMESTVLLAM 128
Db 61 VYFLCMLSTIDLAASVSTVPKLLAIFWCGAGHISASACLAHMFPIHAFCMESTVLLAM 120
QY 129 AFDYVAICHPLRYATILDTTIIAHIGVAAVVRGSLMLPCPFFIGRLNFCQSHVILTY 188
Db 121 AFDYVAICHPLRYATILDTTIIAHIGVAAVVRGSLMLPCPFFIGRLNFCQSHVILTY 180
QY 189 CEHMAVVKLAGDTRPNRVYGLTAALVIGVDLFCIGLSYALIAQAVRLSSHEARSKAL 248
Db 181 CEHMAVVKLAGDTRPNRVYGLTAALVIGVDLFCIGLSYALIAQAVRLSSHEARSKAL 240
QY 249 GTCGSHVCVILISYTPALFSPFTHRGHHVPHIHLANVYLLPPALNPVYGVKTKQ 308
Db 241 GTCGSHVCVILISYTPALFSPFTHRGHHVPHIHLANVYLLPPALNPVYGVKTKQ 300
QY 309 IRKRVVRVFGSGQGMGIKASE 329
Db 301 IRKRVVRVFGSGQGMGIKASE 321
RESULT 11
ABR01673
ID ABR01673 standard; protein; 316 AA.
XX XX

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Db 121 ESTVLLAMAFDRYVAICHPLRYATILTDITIAHIGVAAVVRGSLMLPCPFLIGRLNFCQ 180
QY 181 SHVILHTYCEHMAVVKLAGDTRPNRVYGLTAALLVIGVDLFCIGLSYALIAQAQVRLSS 240
Db 181 SHVILHTYCEHMAVVKLAGDTRPNRVYGLTAALLVIGVDLFCIGLSYALSAQAQVRLSS 240
QY 241 HEARSKALGTGCGSHVVCVILLISYTPALFSTFTHRGHVPVHIHILLANVYLLPPALNPV 300
Db 241 HEARSKALGTGCGSHVVCVILLISYTPALFSTFTHRGHVPVHIHILLANVYLLPPALNPV 300
QY 301 YGVKTKQIRKRVVRVFSQGGMGKASE 329
Db 301 YGVKTKQIRKRVVRVFSQGGMGKASE 329

RESULT 8
ID AAG71674 standard; protein; 321 AA.
XX AAG71674;
XX 30-JUL-2001 (first entry)
DE Human olfactory receptor polypeptide, SEQ ID NO: 1355.
XX Human; olfactory receptor; OR; primary scent determination;
KW secondary scent determination; polypeptide library; odour receptor;
KW scent profile; scent fingerprint; scent representation.
XX Homo sapiens.
XX WO200127158-A2.
XX 19-APR-2001.
XX 06-OCT-2000; 2000WO-US027582.
XX 08-OCT-1999; 99US-0158615P.
XX 24-FEB-2000; 2000US-0184809P.
XX (DIGI-) DIGISCENTS.
XX (YEDA) YEDA RES & DEV CO LTD.
XX Bellenson J, Smith D, Lancet D, Glusman G, Fuchs T, Yanai I;
XX WPI; 2001-290713/30.
XX New polynucleotides which encode polypeptides involved in olfactory
PT sensation for identifying olfactory agonists and antagonists.
XX Claim 11; Page 838-839; 1857pp; English.

CC The present sequence is an olfactory receptor which is encoded by one of
CC a number of novel polynucleotides. The polynucleotides can be used in
CC screening for olfactory agonists and antagonists. The methods allow for
CC the determination of primary scents and the identification of the odour
CC receptors used to detect these primary scents. The methods also enable
CC determination of secondary scents and the identification of combinations
CC of odour receptors that are involved in detecting such secondary scents.
CC This enables the construction of a scent representation (also called a
CC scent fingerprint or scent profile), which may be used to re-create and
CC edit scents. Libraries of olfactory receptors are useful for determining
CC the interaction pattern of a composition with the receptors, and can be
CC used for determining differences in the olfactory faculties of different
CC individuals
XX Sequence 321 AA;

Query Match 96.0%; Score 1649; DB 4; Length 321;
Best Local Similarity 98.8%; Pred. No. 2.8e-173;
Matches 317; Conservative 1; Mismatches 3; Indels 0; Gaps 0;
9 MESPHTDVPVFFLLGIPGLEQFHLMLSLPVCGLGTATVGNITILVVVATEPVVHKP 68

Db 1 MESPHTDVPVFFLLGIPGLEQFHLMLSLPVCGLGTATVGNITILVVVATEPVVHKP 60
QY 69 VYLFCLMSTIDLAASVSTVPKLLAIFWCGAGHISASACLAHMFHAFQWMESTVLLAM 128
Db 61 VYLFCLMSTIDLAASVSTVPKLLAIFWCGAGHISASACLAQWFFHAFQWMESTVLLAM 120
QY 129 AFDYVAICHPLRYATILTDITIAHIGVAAVVRGSLMLPCPFLIGRLNFCQSHVILHTY 188
Db 121 AFDYVAICHPLRYATILTDITIAHIGVAAVVRGSLMLPCPFLIGRLNFCQSHVILHTY 180
QY 189 CEHMAVVKLAGDTRPNRVYGLTAALLVIGVDLFCIGLSYALIAQAQVRLSSHEARSKAL 248
Db 181 CEHMAVVKLAGDTRPNRVYGLTAALLVIGVDLFCIGLSYALSAQAQVRLSSHEARSKAL 240
QY 249 GTCGSHVVCVILLISYTPALFSTFTHRGHVPVHIHILLANVYLLPPALNPVYGVKTKQ 308
Db 241 GTCGSHVVCVILLISYTPALFSTFTHRGHVPVHIHILLANVYLLPPALNPVYGVKTKQ 300
QY 309 IRKRVVRVFSQGGMGKASE 329
Db 301 IRKRVVRVFSQGGMGKASE 321

RESULT 9
ID AAU80511 standard; protein; 321 AA.
XX AAU80511;
XX 12-MAR-2002 (first entry)
DE Human G-coupled receptor (GCREC) protein, Seq ID No 19.
XX Human; cytostatic; neuroprotective; immunosuppressant; nootropic;
KW anti-inflammatory; anti-viral; gastrointestinal; cardiovascular;
KW cerebroprotective; G-coupled receptor; cell proliferative disease;
KW lymphoma; leukaemia; breast cancer; cirrhosis; neurological disorder;
KW stroke; Alzheimer's disease; multiple sclerosis; mental retardation;
KW cardiovascular disease; atherosclerosis; angina pectoris; indigestion;
KW congestive heart failure; gastrointestinal disorder; dysphagia; AIDS;
KW gastritis; autoimmune disorder; inflammatory disorder; Crohn's disease;
KW systemic lupus erythematosus; metabolic disorder; diabetes; obesity;
KW viral infection; herpesvirus; parvovirus;
KW acquired immune deficiency syndrome.

XX Homo sapiens.
XX WO200190359-A2.
XX 29-NOV-2001.
XX 22-MAY-2001; 2001WO-US016833.
XX 22-MAY-2000; 2000US-0206222P.
XX 25-MAY-2000; 2000US-0207476P.
XX 02-JUN-2000; 2000US-0208834P.
XX 07-JUN-2000; 2000US-0208861P.
XX (INCY-) INCYTE GENOMICS INC.
XX Patterson C, Tribouley CM, Yao MG, Griffin JA, Thornton M, Lu Y;
PI Kallick DA, Gandhi AR, Au-Young J;
XX WPI; 2002-106199/14.
XX N-PSDB; ABK16633.
XX DR 2002-106199/14.
XX N-PSDB; ABK16633.
XX New G-protein coupled receptors useful for treating or preventing cell
PT proliferative (e.g. leukemia), neurological (e.g. stroke), cardiovascular
PT or autoimmune/inflammatory disorders.
XX Claim 1; Page 133-134; 148pp; English.

XX	AAU85266	standard; protein; 329 AA.
XX	AAU85266;	
XX	AC	
XX	AC	
XX	08-MAY-2002	(first entry)
XX	G-coupled olfactory receptor #127.	
XX	Human; olfactory G-coupled receptor; sensory perception of odourant;	
XX	odour composition; taste composition.	
XX	Homo sapiens.	
OS	WO200198526-A2.	
XX	27-DEC-2001.	
XX	22-JUN-2001; 2001WO-US020122.	
XX	22-JUN-2000; 2000US-0213812P.	
PR	13-MAR-2001; 2001US-00804291.	
XX	(SENO-) SENOMYX INC.	
PA	Zozulya S, Stryer L;	
XX	WPI; 2002-083330/11.	
DR	N-PSDB; ABK37625.	
XX	Representing sensory perception of one or more odorants for the	
PT	identification and design of tastes and odors comprises providing a	
PT	representative group of n olfactory receptors.	
PT	Claim 1; Page 115; 182pp; English.	
PS	The invention relates to a method of representing sensory perception of	
XX	one or more odorants. The method comprises: (a) providing a	
CC	representative class of n olfactory receptors or ligand binding domains	
CC	(LBD) of these receptors; (b) measuring values X1 to Xn representative of	
CC	at least one activity of one or more odorants selected from: (i) binding	
CC	one or more odorants to the LBD of at least one of the n olfactory	
CC	receptors; (ii) activating at least one of the n olfactory receptors with	
CC	the one or more odorants; and (iii) blocking at least one of the n	
CC	olfactory receptors with the one or more odorants; and (c) generating a	
CC	representation of the sensory perception from the values X1 to Xn. The	
CC	representation of the sensory perception of odorants is useful for the	
CC	design and formulation of odour and taste compositions. AAU85140-AAU85393	
CC	represent human olfactory G-coupled receptor amino acid sequences of the	
CC	invention	
XX	Sequence 329 AA;	
XX	SQ	
XX	Query Match	98.4%; Score 1691; DB 5; Length 329;
XX	Best Local Similarity	98.8%; Pred. No. 6.7e-178;
XX	Matches 325; Conservative	1; Mismatches 3; Indels 0; Gaps 0;
QY	1	MSSTLGHNMESPHHTDVPSPVFFLLGIPGLEQFHLMSLPVCGLGCTATVGNITILVVVA 60
DB	1	MSSTLGHNMESPHHTDVPSPVFFLLGIPGLEQFHLMSLPVCGLGCTATVGNITILVVVA 60
QY	61	TEPVLHKPVYLFCLMLSTIDLAAASVSTVPKLLAIFWCAGHISASACLAHMFHAFQCM 120
DB	61	TEPVLHKPVYLFCLMLSTIDLAAASVSTVPKLLAIFWCAGHISASACLAHMFHAFQCM 120
QY	121	ESTVLLAMAFDRYVAICHPLRYATILDTDTIAHIGVAAVVRGSLMLPCPPFFIGRLNFCQ 180
DB	121	ESTVLLAMAFDRYVAICHPLRYATILDTDTIAHIGVAAVVRGSLMLPCPPFFIGRLNFCQ 180
QY	181	SHVILHTYCEHMAVVKLACGDTDRNRYVGLTAALLVIVGDVDFCTGLSVALTAQAVRLUSS 240

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Page 8

CC polynucleotide sequences shown in ADC85548-ADC87616 encode GPCR's of the
CC invention.

XX SQ Sequence 1390 BP; 317 A; 353 C; 293 G; 427 T; 0 U; 0 Other;

Alignment Scores:
Pred. No.: 8 03e-169 Length: 1390
Score: 1691.00 Matches: 325
Percent Similarity: 99.09% Conservatives: 1
Best Local Similarity: 98.78% Mismatches: 3
Query Match: 98.43% Indels: 0
DB: 9 Gaps: 0

US-10-081-775-2 (1-329) x ADC86344 (1-1390)

Qy 1 MetSerThrLeuGlyHisAsnMetGluSerProHisHisThrAspValAspProSer 20
Db 201 ATGTCAGCACTCTGGCCACAAATGCTCTTAATCACTGATGTTGACCTTCT 260
Qy 21 ValPhePheLeuLeuGlyLeuProGlyLeuGluGlnPheHisLeuTrpLeuSerLeuPro 40
Db 261 GTCATCTCTCCCTGGGATCCAGGCTGGAACAATTTCAATTTGTGGCTCTCACTCCCT 320
Qy 41 ValCysGlyLeuGlyThrAlaThrIleValGlyAsnIleThrIleLeuValValAla 60
Db 321 GTGTGGCTTAGGCACAGCCACAAATGTCGGCATATACTATTTCTGGTTGTGTGCC 380
Qy 61 ThrGluProValLeuHisIleProValTyrLeuPheLeuCysMetLeuSerThrIleAsp 80
Db 381 ACTGAACAGTCTTGCCACAGCTGTGTACCTTTTCTGTGCATGCTCTCAACCATCGAC 440
Qy 81 LeuAlaAlaSerValSerThrValProIleLeuAlaIlePheTrpCysGlyAlaGly 100
Db 441 TTGCTGCTCTGTCTCCACAGTCTCCCAAGCTACTGGCTATCTTCTGGTGGAGCCGGA 500
Qy 101 HisIleSerAlaSerAlaCysLeuAlaHisMetPhePheIleHisAlaPheCysMetMet 120
Db 501 CATATATCTGCTCTGCTGCTGGCAGATGTTCTTCATTCATGCTTCTGATGATG 560
Qy 121 GluSerThrValLeuLeuAlaMetAlaPheAspArgTyrValAlaIleCysHisProLeu 140
Db 561 GAGTCCACTGTGCTACTGGCCATGCGCTTGTATGCTAGTGGCCATCTGCCACCCACTC 620
Qy 141 ArgTyrAlaThrIleLeuThrAspThrIleAlaHisIleGlyValAlaAlaVal 160
Db 621 CGCTATGCCAATCTCTCAGTACACCATCATGTCCTCCACATAGGCGGTGGAGCTGTAGT 680
Qy 161 ArgGlySerLeuLeuMetLeuProCysProPhePheIleGlyArgLeuAsnPheCysGln 180
Db 681 CGAGGCTCCCTGCTCATGCTCCCATGCTCCCTTCTTATTTGGCGTTGAACTTCTGCCNA 740
Qy 181 SerHisValIleLeuHisThrTyrCysGluHisMetAlaValValLeuAlaCysGly 200
Db 741 AGCATGTGATCTTACACACCTACTGTGAGCACATGCTGTGTGTGAGTGGCTGTGGA 800
Qy 201 AspThrArgProAsnArgValTyrGlyLeuThrAlaLeuLeuValIleGlyValAsp 220
Db 801 GACACAGGCTACCCGTGTGTGGGTGACAGCTGCACGTGTGGTTCATTTGGGGTTGAC 860
Qy 221 LeuPheCysIleGlyLeuSerTyrAlaLeuIleAlaGlnAlaValLeuLeuSerSer 240
Db 861 TTGTTTGTGATGCTCTCTCTATGCTTGTAGTGCACNAGTGTCTTCTGCTCTCTATCC 920
Qy 241 HisGluAlaArgSerLysAlaLeuGlyThrCysGlySerHisValCysValIleLeuIle 260
Db 921 CATGAAGCTCGTCCAGGCTTGGGACCTGTGGTTCCTCATGCTGTGTGCTCATCTCATC 980
Qy 261 SerTyrThrProAlaLeuPheSerPhePheThrHisArgPheGlyHisHisValProVal 280
Db 981 TCTTATACACAGGCTCTCTCTCTCTTTTACACACCGCTTGGCCATCATCAGTCTCCAGTC 1040
Qy 281 HisIleHisIleLeuLeuAlaAsnValTyrLeuLeuLeuProAlaLeuAsnProVal 300

Db 1041 CATATTACATCTTTTGGCCAAATGTTATCTGCTTTTGCACCTGCTCTTAATCTGTG 1100

Qy 301 ValTyrGlyValLysThrLysGlnIleArgLysArgValValArgValPheGlnSerGly 320

Db 1101 GTATATGGAGTTAAGACCAACACAGATCCGTAAGAGAGTGTTCAGGGTGTTCAGAGTGG 1160

Qy 321 GlnGlyMetGlyIleLysAlaSerGlu 329

Db 1161 CAGGAAATGGGCATCAAGGCATCTGAG 1187

RESULT 8

AAH31850

ID AAH31850 standard; DNA; 963 BP.

XX AAH31850;

DT 30-JUL-2001 (first entry)

XX Human olfactory receptor polynucleotide, SEQ ID NO: 423.

XX Human; olfactory receptor; OR; primary scent determination;

XX secondary scent determination; polypeptide library; odour receptor;

XX scent profile; scent fingerprint; scent representation; ds.

XX Homo sapiens.

XX WO200127158-A2.

XX 19-APR-2001.

XX 06-OCT-2000; 2000WO-US027582.

XX 08-OCT-1999; 99US-0158615P.

XX 24-FEB-2000; 2000US-0184809P.

XX (DIGI-) DIGISCENTS.

XX (YEDA) YEDA RES & DEV CO LTD.

XX Bellenson J, Smith D, Lancet D, Glusman G, Fuchs T, Yanai I;

XX WPI; 2001-290713/30.

XX New polynucleotides which encode polypeptides involved in olfactory

XX sensation for identifying olfactory agonists and antagonists.

XX Claim 8; Page 349; 1857pp; English.

XX The present sequence is one of a number of isolated polynucleotides which

XX encode polypeptides involved in olfactory sensation. The polynucleotides

XX can be used in screening for olfactory agonists and antagonists. The

XX methods allow for the determination of primary scents and the

XX identification of the odour receptors used to detect these primary

XX scents. The methods also enable determination of secondary scents and the

XX identification of combinations of odour receptors that are involved in

XX detecting such secondary scents. This enables the construction of a scent

XX representation (also called a scent fingerprint or scent profile), which

XX may be used to re-create and edit scents. Libraries of olfactory

XX receptors are useful for determining the interaction pattern of a

XX composition with the receptors, and can be used for determining

XX differences in the olfactory faculties of different individuals

SQ Sequence 963 BP; 181 A; 276 C; 216 G; 290 T; 0 U; 0 Other;

Alignment Scores:

Pred. No.: 1 13e-164 Length: 963
Score: 1649.00 Matches: 317
Percent Similarity: 99.07% Conservatives: 1
Best Local Similarity: 98.75% Mismatches: 3
Query Match: 95.98% Indels: 0
DB: 4 Gaps: 0

US-10-081-775-2 (1-329) x AAH31850 (1-963)

QY 9 MetGluSerProHisHisThrAspValAspProSerValPhePheLeuLeuGlyIlePro 28
DB 1 ATGGAATCTCTAATACACTGATGTGACCCCTTCTGCTTCTTCTCTCTGGGCAATCCCA 60
QY 29 GlyLeuGluGlnPheHisLeuTrpLeuSerLeuProValCysGlyLeuGlyThrAlaThr 48
DB 61 GGTCTGGAACAATTTCAATTTGGCTCTCACTCCCTGTGTGTGGCTTAGGCACAGCCACA 120
QY 49 IleValGlyAsnIleThrIleValValAlaValAlaThrGluProValLeuHisLysPro 68
DB 121 ATGTGGGGCAATATACTATTCTGGTGTGTGGCACTGAACCACTCTTGCAACAGCT 180
QY 69 ValTyrLeuPheLeuCysMetLeuSerThrIleAspLeuAlaAlaSerValSerThrVal 88
DB 181 GTGTACCTTTTCTGTGATGCTCTCAACATCGATTGGTGGCTCTGTCTCCAGATT 240
QY 89 ProLysLeuLeuAlaIlePheTrpCysGlyAlaGlyHisIleSerAlaSerAlaCysLeu 108
DB 241 CCCAAGCTACTGGCTATCTTCTGGTGTGGAGCGGACATATATCTGCTCTGCTGCTG 300
QY 109 AlaHisMetPhePheIleHisAlaPheCysMetMetGluSerThrValLeuLeuAlaMet 128
DB 301 GCACAGATGTTCTTCAATTCATGCTTCTCATGATGGAGTCCACTGCTGCTACTGGCCATG 360
QY 129 AlaPheAspArgTyrValAlaIleCysHisProLeuArgTyrAlaThrIleLeuThrAsp 148
DB 361 GCCTTTGATCGTAGTGGCCATCTGCCACCCACTCCGCTATGCCAATCTCTCATCTGAC 420
QY 149 ThrIleIleAlaHisIleGlyValAlaAlaValAlaValArgLysLeuLeuMetLeuPro 168
DB 421 ACCATCATTTGCCACATAGGGGTGGCAGCTGTAGTGGAGGCTCCCTGCTCATGCTCCCA 480
QY 169 CysProPhePheIleGlyArgLeuAsnPheCysGlnSerHisValIleLeuHisThrTyr 188
DB 481 TGTCCCTCTCTTATTGGGGCTTTGAATCTCTGCCAAGCCATGTGATCTCTACACAGTAC 540
QY 189 CysGluHisMetAlaValValLysLeuAlaCysGlyAspThrArgProAsnArgValTyr 208
DB 541 TGTGAGCATGCTGCTGTGGAGCTGGCTGTGGAGACACAGGCTTAACCGTGTGTAT 600
QY 209 GlyLeuThrAlaAlaLeuLeuValIleGlyValAspLeuPheCysIleGlyLeuSerTyr 228
DB 601 GGGCTGACAGCTGCACTGTGTGCTTGGGTTGACTTGTGTTCATTTGCTCTCTCTAT 660
QY 229 AlaLeuIleAlaGlnAlaValLeuArgLeuSerSerHisGluAlaArgSerLysAlaLeu 248
DB 661 GCCCTAAGTGCAAGCTGCTCTGCTGCTCTATCCATGCAAGCTGCTGCAAGGCCCTA 720
QY 249 GlyThrCysGlySerHisValCysValIleLeuIleSerTyrThrProAlaLeuPheSer 268
DB 721 GGGAGCTGTGTTCCTCATGCTGTGTCATCTCTTATACACAGCCCTCTTCTCC 780
QY 269 PhePheThrHisArgPheGlyHisHisValProValHisIleHisIleLeuLeuAlaAsn 288
DB 781 TTTTATACACACCGCTTGGCCATCATCGTTCAGTCCATATTCACATCTTTTGGCCAT 840
QY 289 ValTyrLeuLeuLeuProAlaLeuAsnProValValTyrGlyValLysThrLysGln 308
DB 841 GTTATCTGCTTTTGGACCTGCTCTTAATCTCTGTGTATGAGTTAAGACCAACAG 900
QY 309 IleArgLysArgValValArgValPheGlnSerGlyGlnGlyMetGlyIleLysAlaSer 328
DB 901 ATCCGTAAAGAGTTGTGAGGGTGTTCAGAGTGGGCAAGGGAATGGGCAATCAAGGCATCT 960
QY 329 Glu 329
DB 961 GAG 963

RESULT 9
ID ABK16633
XX ABK16633 standard; cDNA; 966 BP.
AC ABK16633;

XX 14-MAR-2002 (first entry)
DT Human G-coupled receptor (GCRC) cDNA, Seq ID No 42.
XX Human; cytostatic; neuroprotective; immunosuppressant; nootropic;
XX anti-inflammatory; anti-viral; gastrointestinal; cardiovascular;
XX cerebroprotective; G-coupled receptor; cell proliferative disease;
XX lymphoma; leukemia; breast cancer; cirrhosis; neurological disorder;
XX stroke; Alzheimer's disease; multiple sclerosis; mental retardation;
XX cardiovascular disease; atherosclerosis; angina pectoris; indigestion;
XX congestive heart failure; gastrointestinal disorder; dysphagia; AIDS;
XX gastritis; autoimmune disorder; inflammatory disorder; Crohn's disease;
XX systemic lupus erythematosus; metabolic disorder; diabetes; obesity;
XX viral infection; herpesvirus; parvovirus;
XX acquired immune deficiency syndrome; ss.
XX Homo sapiens.
OS WO200190359-A2.
XX 29-NOV-2001.
XX 22-MAY-2001; 2001WO-US016833.
XX 22-MAY-2000; 2000US-0206222P.
PR 25-MAY-2000; 2000US-0207476P.
PR 02-JUN-2000; 2000US-0208834P.
PR 02-JUN-2000; 2000US-0208861P.
PR 07-JUN-2000; 2000US-0209868P.
XX (INCY-) INCYTE GENOMICS INC.
XX Patterson C, Tribouley CM, Yao MG, Griffin JA, Thornton M, Lu Y;
PI Kallick DA, Gandhi AR, Au-Young J;
XX WPI; 2002-106199/14.
DR P-PSDB; AA080511.
DR New G-protein coupled receptors useful for treating or preventing cell
XX proliferative (e.g. leukemia), neurological (e.g. stroke), cardiovascular
or autoimmune/inflammatory disorders.
PS Claim 5; Page 146; 148pp; English.
XX The invention relates to a novel human G-coupled receptor (I). (I) and
CC its corresponding polynucleotides are useful for diagnosing, treating or
CC preventing cell proliferative diseases (e.g. lymphoma, leukemia, breast
CC cancer or cirrhosis), neurological disorders (e.g. stroke, Alzheimer's
CC disease, multiple sclerosis or mental retardation), cardiovascular
CC diseases (e.g. atherosclerosis, angina pectoris or congestive heart
CC failure), gastrointestinal disorders (e.g. dysphagia, indigestion or
CC gastritis), autoimmune/inflammatory disorders (e.g. AIDS, Crohn's disease
CC or systemic lupus erythematosus) or metabolic disorders (e.g. diabetes or
CC obesity), or viral infections (e.g. infection by herpesvirus or
CC parvovirus). ABK16633-ABK16637 represent novel human G-coupled receptor
CC coding sequences of the invention
XX Sequence 966 BP; 182 A; 276 C; 217 G; 291 T; 0 U; 0 Other;

Alignment Scores:
Pred. No.: 1.34e-164 Length: 966
Score: 1649.00 Matches: 317
Percent Similarity: 99.07% Conservative: 1
Best Local Similarity: 98.75% Mismatches: 3
Query Match: 95.98% Indels: 0
DB: Gaps: 0

US-10-081-775-2 (1-329) x ABK16633 (1-966)

QY 9 MetGluSerProHisHisThrAspValAspProSerValPhePheLeuLeuGlyIlePro 28
DB 1 ATGGAATCTCTAATACACTGATGTGACCCCTTCTGCTTCTTCTCTCTGGGCAATCCCA 60

Human DNA for olfactory and pheromone G protein-coupled receptor #212..

29	QY	GlyLeuGluGlnPheHisLeuTrpLeuSerLeuProValCysGlyLeuGlyThrAlaThr	48
61	DB	GGCTGGAAACAATTCATTGTGGCTCTCACTCCCTGTGTGGCTAGGCACAGCCACA	120
49	QY	IleValGlyAsnIleThrIleLeuValValAlaThrGluProValLeuHisIlePro	68
121	DB	ATTGTGGGCATATAAATATTCTGGTTGTGTGCCACTGAACCAAGTCTTGACAAAGCT	180
69	QY	ValTrpLeuPheLeuCysMetLeuSerThrIleAspLeuAlaAlaSerValSerThrVal	88
181	DB	GTGTACCTTTTCTGTGTCATGCTCTCAACCAATCGACTTGGCTGGCTCTGTCTCCACAGTT	240
89	QY	ProLysLeuLeuAlaIlePheTrpCysGlyAlaGlyHisIleSerAlaSerAlaCysLeu	108
241	DB	CCCAGCTACTGGCTATCTCTGGTGTGGAGCGGACATATATCTGCCTCTGCCTGCCTG	300
109	QY	AlaHisMetPhePheIleHisAlaPheCysMetMetGluSerThrValLeuLeuAlaMet	128
301	DB	GCACAGATGTTCTTCAATTCATGGCTTCTGCATGATGGAGTCCACTGTGTACTATGGCCATG	360
129	QY	AlaPheAspArgTrpValAlaIleCysHisProLeuArgTrpAlaThrIleLeuThrAsp	148
361	DB	GCCTTTGATCGCTAGTGGCCATCTGCCACCACCTCCGCTATGCCACAATCTCTACTGAC	420
149	QY	ThrIleAlaHisIleGlyValAlaAlaValAlaArgGlySerLeuLeuMetLeuPro	168
421	DB	ACCATCATTTGCCACATAGCGGTGGCAGCTGTAGTGCAGAGGTCCCTGTCTATGTCTCCA	480
169	QY	CysProPhePheIleGlyArgIleuAsnPheCysGlnSerHisValIleLeuHisThrTrp	188
481	DB	TGTCCCTCTCTTATTGGGGCTTTGAACTCTCTGCCAAAGCCATGTGAATCTTACACAGTAC	540
189	QY	CysGluHisMetAlaValValLysLeuAlaCysGlyAspThrArgProAsnArgValTrp	208
541	DB	TGTGAGCACATGGCTGTGTGTGAAGCTGGCTGTGGAGACACCCAGGCTTAACCGTGTGTAT	600
209	QY	GlyLeuThrAlaAlaLeuLeuValIleGlyValAspLeuPheCysIleGlyLeuSerTrp	228
601	DB	GGCTGCACAGCTGCATGTTGGTTCATTTGGGGTTGACTTGTTTTGCATTTGTCTCTCCTAT	660
229	QY	AlaLeuIleAlaGlnAlaValLeuArgLeuSerSerHisGluAlaArgSerLysAlaLeu	248
661	DB	GGCTTAAGTGCACAGCTGCTTCGCCCTCTCATCCCAAGAGCTGGTCCAGAGGCCCTA	720
249	QY	GlyThrCysGlySerHisValCysValIleLeuIleSerTrpThrProAlaLeuPheSer	268
721	DB	GGGACCTGTGGTTCCTCATGCTGTGCATCCTCATCTCTTATACACAGGCCCTCTCTCC	780
269	QY	PhePheThrHisArgPheGlyHisHisValProValHisIleHisIleLeuLeuAlaAsn	288
781	DB	TTTTTTTACACACCGCTTTGGCCATCAGTTCCAGTCCATATTACATTTCTTTTGGCCAAAT	840
289	QY	ValTrpLeuLeuLeuProProAlaLeuAsnProValValTrpGlyValLysThrLysGln	308
841	DB	GTTTATCTGCTTTTGGCACCTGCTCTTAATCCTGTGTATATATGGAGTTTAAAGCCAAACAG	900
309	QY	IleArgLysArgValValArgValPheGlnSerGlyGlnGlyMetGlyIleLysAlaSer	328
901	DB	ATCTCTTAATAGCTTGTCTAAGTCTTTCAAGTGTGCAGAGGAATGGCATCAAGGCATCT	960

10. T. J. P. 10

RESULT IN
ABK68612

ABK68612 standard; DNA; 966 BP.

AA
AC
ABK68612:

02-III-2002 (first entry)

SECRET
7007-1000-20
X

10

10

1

[illegible]

Alignment Scores:		
Pred. No.:	1,348-164	966
Score:	1649.00	217
Percent Similarity:	99.07%	Conservative:
Best Local Similarity:	98.75%	Mismatches:
Query Match:	95.98%	Indels:
DB:	5	Gaps:
DB:		

US-10-081-775-2 (1-329) x ARK68612 (1-966)

9 MetGluSerProHisHisThrAspValAspProSerValPheLeuGlyIlePro 28
:::
1 MetGluSerProHisHisThrAspValAspProSerValPheLeuGlyIlePro 60

